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TRANSMITTAL FORM		Application Number	09/463,470	-RECE	IVED
		Filing Date	January 21, 2	.000	
		First Named Inventor	Morten Soega	aard AUG 2	P 2001
(to be used for all correspondence after initial filing)		Group Art Unit	N/A	TECH CENTER	1600/2900
(0.00.00,00,00,00,00,00,00,00,00,00,00,00		Examiner Name	Not Yet Assig	ned	
Total Number of Pages in This Submission		Attorney Docket Numb	er HO-P01938U	S0	
ENCLOSURES (check all that apply)					
X Fee Transmittal Form	Assignment I		After Allowance to Group	Communication	
Fee Attached	Drawing(s)		Appeal Commu Appeals and Int	nication to Board of erferences	
Amendment/Reply	Licensing-rel	ated Papers	Appeal Commun (Appeal Notice, Brid	ication to Group	
After Final	Petition		Proprietary Info	mation	
Affidavits/declaration(s)	Petition to Convert to a Provisional Status Lette		Status Letter		
Extension of Time Request	Power of Attorney, Revocation Change of Correspondence Address X Other Enclosure(s) (please identify below)			e(s) ow)	
Express Abandonment Request	Terminal Disclaimer Sequence Listing Transmittal Sequence Listing				
Information Disclosure Statement	Request for	Refund	Diskette (1 copy) Copy of Notice to		
Certified Copy of Priority Document(s)	CD, Number	of CD(s)	Return Post Card		
Response to Missing Parts/ Incomplete Application	Remarks				
Response to Missing Parts under 37 CFR 1.52 or 1.53					
under 67 61 10 1.02 61 1.06					
SIGNATURE OF APPLICANT, ATTORNEY, OR AGENT					1
Firm or Individual Name FULBRIGHT & JAWORSKI L.L.P. Melissa W. Acosta					
Signature					
Date August 23, 2001					

Transmittal

I hereby certify that this correspondence is being deposited with the U.S. Postal Service as Express Mail, Airbill No. EK102656123US, in an envelope addressed to: Commissioner for Patents, Washington, DC 20231, on the date shown below. Signature:

Dated: August 23, 2001

Application No.: 09/163470

NOTICE TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE DISCLOSURES

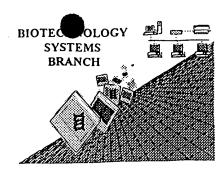
Applicant must file the items indicated below within the time period set the Office action to which the Notice is attached to avoid abandonment under 35 U.S.C. § 133 (extensions of time may be obtained under the provisions of 37 CFR 1.136(a)).

The nucleotide and/or amino acid sequence disclosure contained in this application does not comply with the requirements for such a disclosure as set forth in 37 C.F.R. 1.821 - 1.825 for the following reason(s):

Ä	attention is directed to the final rulemaking notice published at 55 FR 18230 (May 1, 1990), and 1114 OG 29 (May 15, 1990). If the effective filing date is on or after July 1, 1998, see the final rulemaking notice published at 63 FR 29620 (June 1, 1998) and 1211 OG 82 (June 23, 1998).
	2. This application does not contain, as a separate part of the disclosure on paper copy, a "Sequence Listing" as required by 37 C.F.R. 1.821(c).
	3. A copy of the "Sequence Listing" in computer readable form has not been submitted as required by 37 C.F.R. 1.821(e).
X	4. A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 C.F.R. 1.822 and/or 1.823, as indicated on the attached copy of the marked -up "Raw Sequence Listing."
	5. The computer readable form that has been filed with this application has been found to be damaged and/or unreadable as indicated on the attached CRF Diskette Problem Report. A Substitute computer readable form must be submitted as required by 37 C.F.R. 1.825(d).
	6. The paper copy of the "Sequence Listing" is not the same as the computer readable from of the "Sequence Listing" as required by 37 C.F.R. 1.821(e).
	7. Other:
Аp	plicant Must Provide:
V	An initial or substitute computer readable form (CRF) copy of the "Sequence Listing".
×	Corresponding An initial of substitute paper copy of the "Sequence Listing", as well as an amendment directing its entry into the specification.
K	A statement that the content of the paper and computer readable copies are the same and, where applicable, include no new matter, as required by 37 C.F.R. 1.821(e) or 1.821(f) or 1.821(g) or 1.825(b) or 1.825(d).
Fo	r questions regarding compliance to these requirements, please contact:
For	r Rules Interpretation, call (703) 308-4216 r CRF Submission Help, call (703) 308-4212 tentin Software Program Support
	Technical Assistance703-287-0200 To Purchase Patentin Software703-306-2600
	PLEASE RETURN A COPY OF THIS NOTICE WITH YOUR REPLY



RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	<u>09/463,470 A</u>
Source:	1644
Date Processed by STIC:	3-14-01

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address: http://www.uspto.gov/web/offices/pac/checker



1644

RAW SECHENCE LISTING DATE: 03/14/2001 TIME: 10:20:18 PATENT APPLICATION: US/09/463,470A

Input Set : A:\es.txt

Output Set: N:\CRF3\03142001\I463470A.raw

Does Not Comply Corrected Diskette Needed 3 <110> APPLICANT: SOEGAARD, MORTEN ABRAHMSEN, LARS 4 pp. 1,2 LANDO, PETER 5 6 FORSBERG, GORAN 7 KALLAND, TERJE DOHLSTEN, MIKAEL 10 <120> TITLE OF INVENTION: CYTOLYSIS OF TARGET CELLS BY SUPERANTIGEN CONJUGATES INDUCING T-CELL 11 ACTIVATION 13 <130> FILE REFERENCE: P01938US0; 10001907 15 <140> CURRENT APPLICATION NUMBER: 09/463,470A 16 <141> CURRENT FILING DATE: 2000-01-21 18 <150> PRIOR APPLICATION NUMBER: 60/053,211 19 <151> PRIOR FILING DATE: 1997-07-21 21 <150> PRIOR APPLICATION NUMBER: PCT/EP98/04219 22 <151> PRIOR FILING DATE: 1998-07-21 24 <150> PRIOR APPLICATION NUMBER: 9704170-1 25 <151> PRIOR FILING DATE: 1997-11-14 27 <160> NUMBER OF SEQ ID NOS: 23 29 <170> SOFTWARE: PatentIn version 3.0

ERRORED SEQUENCES

386 <210> SEQ ID NO: 23 387 <211> LENGTH: 17 388 <212> TYPE: PRT 389 <213> ORGANISM: ARTIFICIAL SEQUENCE 391 <220> FEATURE: 392 <221> NAME/KEY: misc_feature 393 <222> LOCATION: (1)..(17) 394 <223> OTHER INFORMATION: Designated peptide to act as a Q-linker. 397 <400> SEQUENCE: 23 399 Gly Pro Arg Gln Ala Asn Glu Leu Pro Gly Ala Pro Ser Gln Glu Glu 400 1 402 Arg 405 25001022.1 E--> 406 7 410 25001022.1 E--> 411 2

> Delete extraneous material from the end of the file. It is causing an invalid amino acid count the sequence # 23. See # 2
 NB 23 200 6
 MB 23

Arg



VERIFICATION SUMMARY

PATENT APPLICATION: US/09/463,470A

DATE: 03/14/2001 TIME: 10:20:19

Input Set : A:\es.txt

Output Set: N:\CRF3\03142001\I463470A.raw

L:359 M:258 W: Mandatory Feature missing, <220> FEATURE:

L:359 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION: L:406 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:23 M:332 Repeated in SeqNo=23